

↓

A)	<u>MRGATRVSIMLLLVTVSDC</u>	AVITGA
B)	<u>MRSLCCAPLLLLLLLPPLLLTPRAGDA</u>	AVITGA
C)	<u>MKCFAQIVVLLLVIAFSHG</u>	AVITGA
D)		AVITGA

CERDVQCGAGTCCAI~~S~~LWLRGLRMCTPLGREGE~~E~~CHPG
CDKDSQCGGGMCCAVSIWVKSIRICTPMGKLGD~~S~~CHPL
CDKDVQCGSGTCCAASAWSRNIRFCIPLGNSGEDCHPA
CERDLQCGKGTCCAVSLWIKSVRVCTPVGTSGEDCHPA
* * ** * *

SHKVPFFRKRKHHTCPCLPNLLCSRFPDGRYRCSMDLKNINF
TRKVPFFGRRMHHTCPCLPGLACLRTSFNRFICLAQK
SHKVPYDGKRLSSLCPCCKSGLTCSK.SGEKFKCS
SHKIPFSGQRMHHTCPCAPNLACVQTSPKKFKCLSKS
* * * *

FIGURE 1

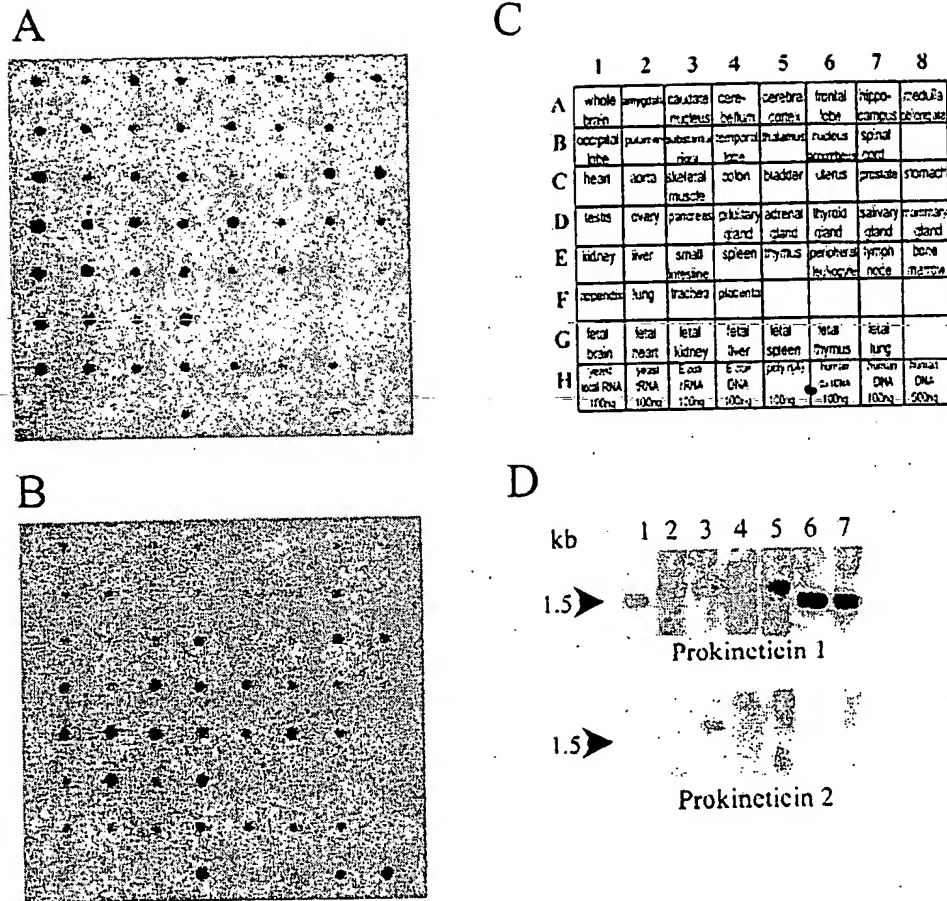


FIGURE 2

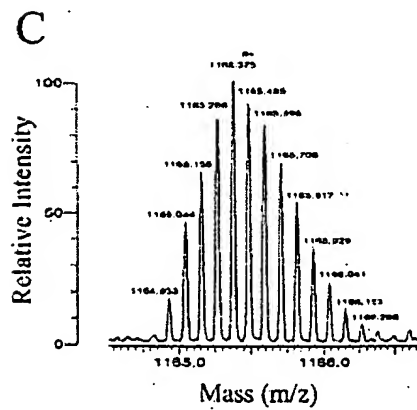
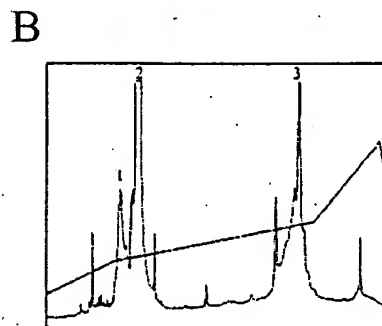
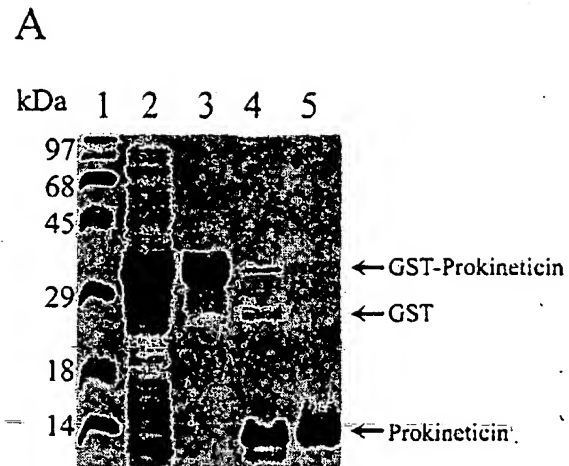


FIGURE 3

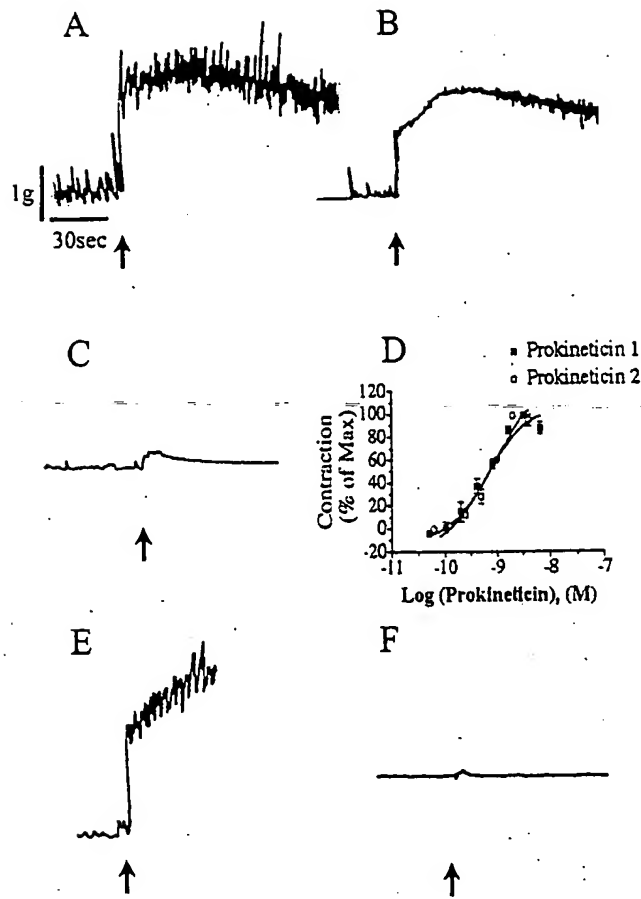
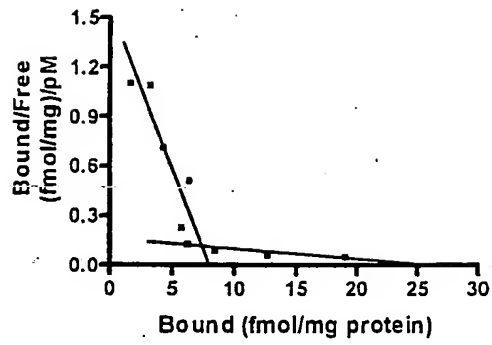


FIGURE 4

A



B

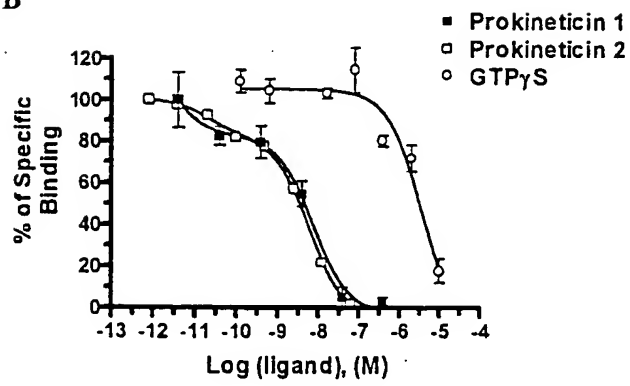


FIGURE 5

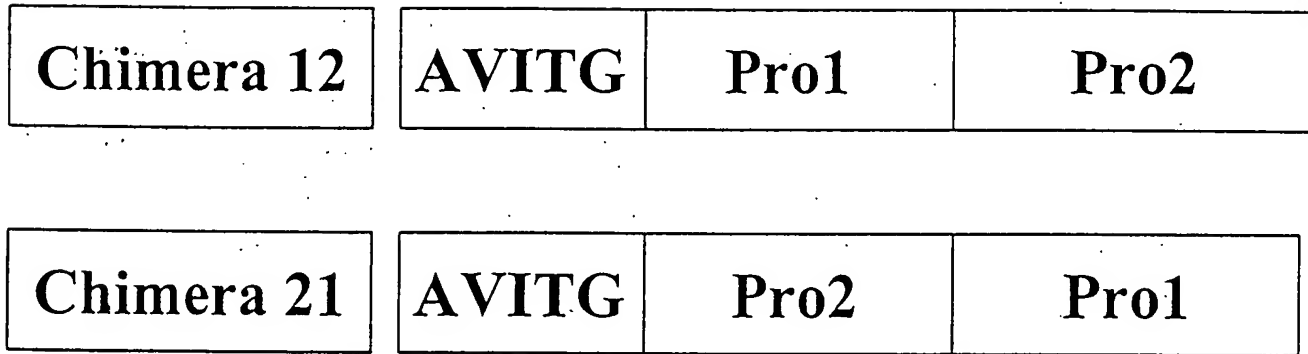
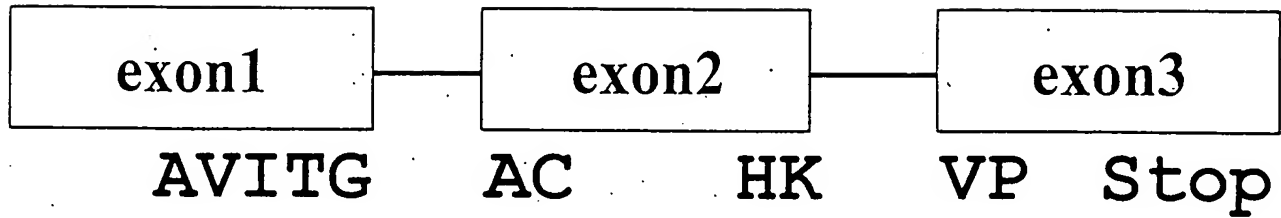
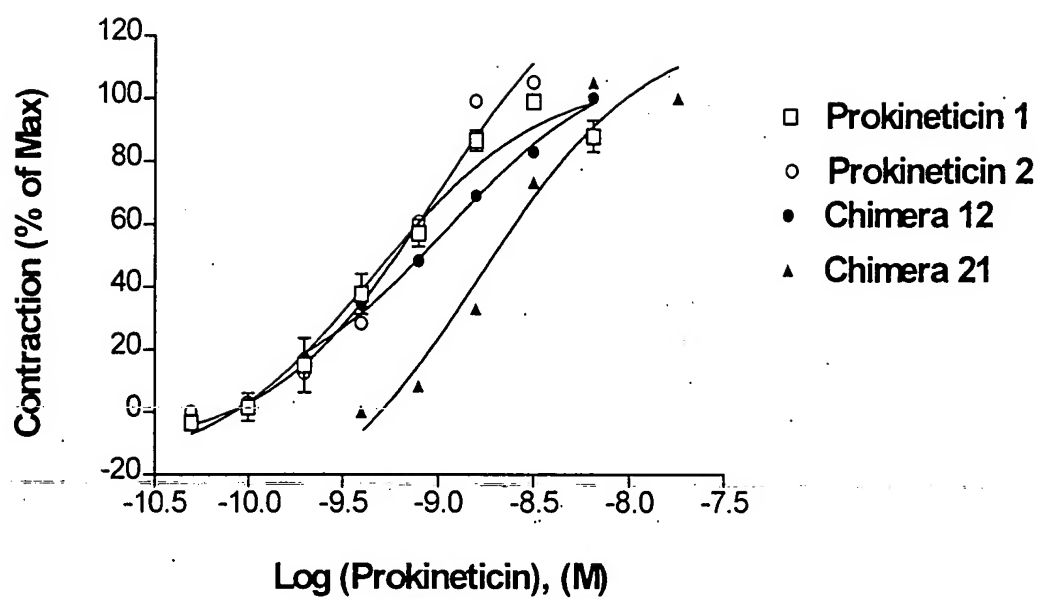


FIGURE 6

A



B

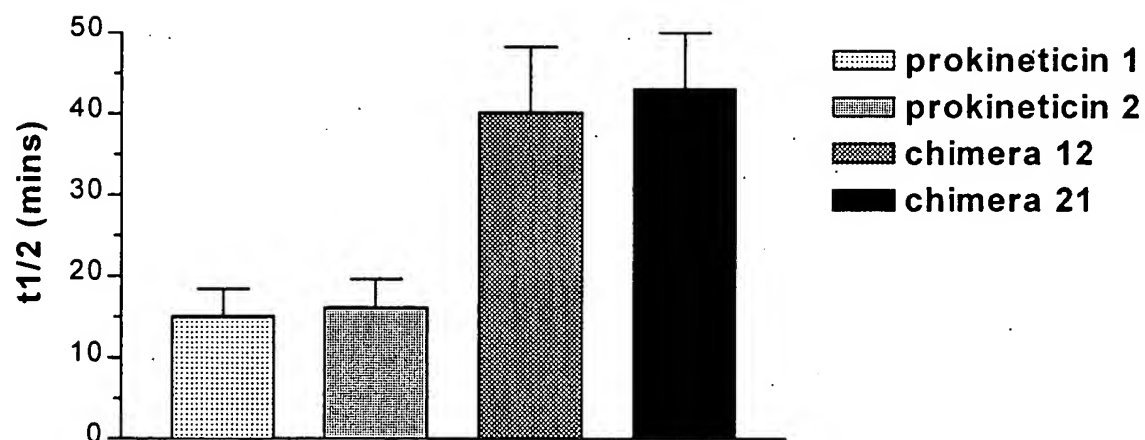
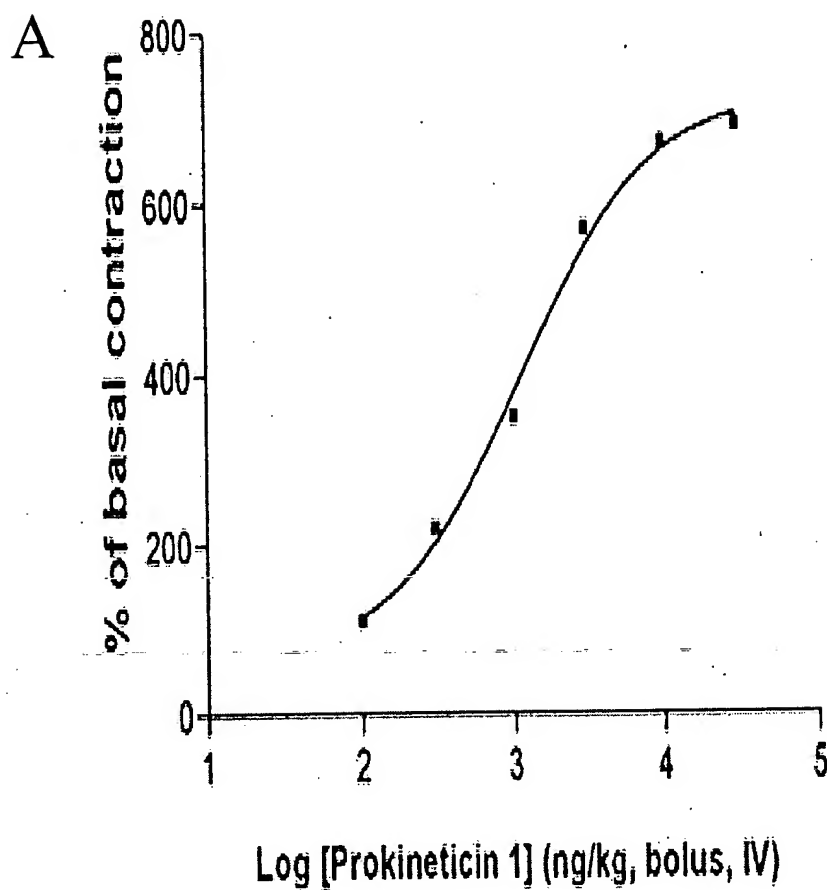


FIGURE 7



B

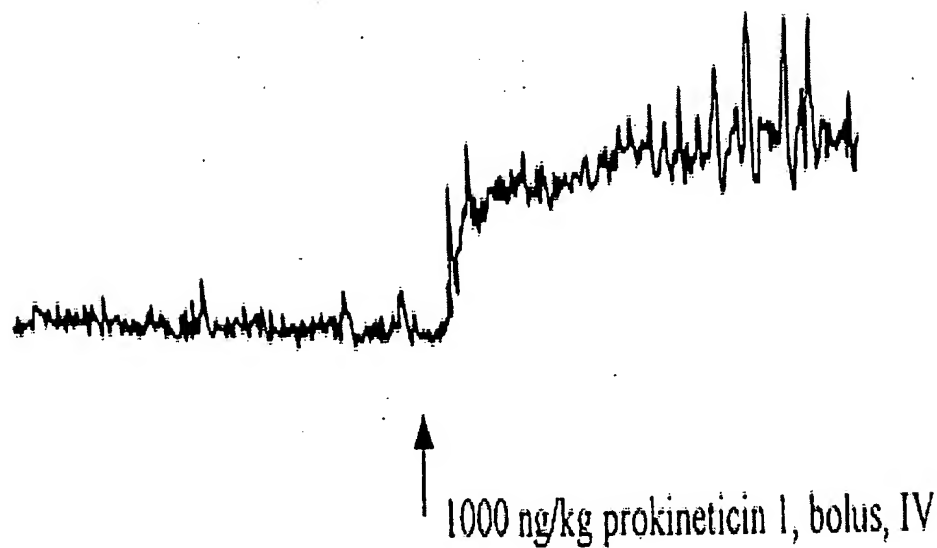


FIGURE 8

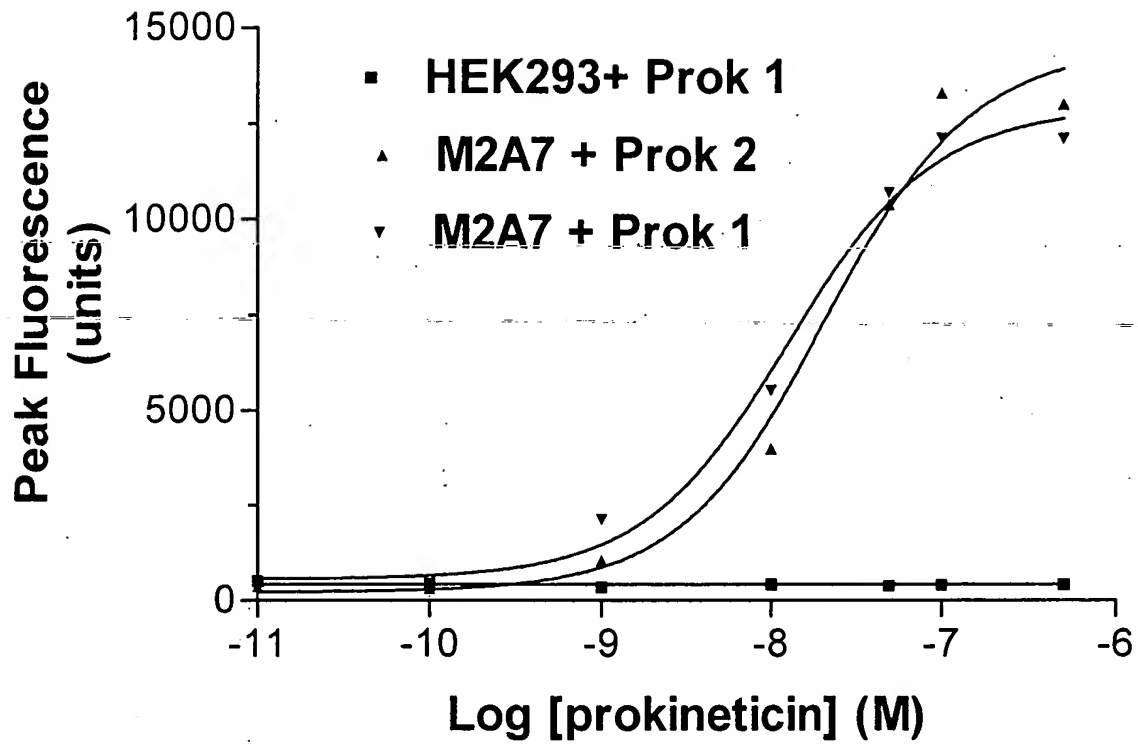


FIGURE 9